



SEQUENCE LISTING

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<120> HEAT SHOCK PROTEIN-BASED VACCINES AND
IMMUNOTHERAPIES

<130> 8449-405-999

<140> 10/776,521

<141> 2004-02-12

<150> 60/503,417

<151> 2003-09-16

<150> 60/463,746

<151> 2003-04-18

<150> 60/462,469

<151> 2003-04-11

<150> 60/447,142

<151> 2003-02-13

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<222> 1, 3, 5, 7

<223> Xaa = hydrophobic amino acid residues

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<223> motif in heptameric region recognized by heat shock protein

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<222> 4, 6

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<213> Lymphocytic Choriomeningitis Virus (LCMV)

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Trp Leu Ser Leu Leu Val Pro Phe Val
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Gly Leu Ser Pro Thr Val Trp Leu Ser Val
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Leu Met Gly Tyr Ile Pro Leu Val Gly Ala
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Ile Leu Lys Glu Pro Val His Gly Val
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Val Glu Ala Glu Ile Ala His Gln Ile
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 <213> *E.coli*

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 <213> *P. falciparum*

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 Lys Ser Lys Asp Glu Leu Asp Tyr
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Trp residue

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Trp residue

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Trp residue

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Trp residue

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Trp residue

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<223> Heat shock protein binding domain with a terminal
Trp residue

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<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 138

Trp Gly Pro Trp Ser Phe Gly Trp
1 5

<210> 139
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 139
 Ala Ile Pro Gly Lys Val Gly Trp
 1 5

 <210> 140
 <211> 8
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 140
 Arg Val His Asp Pro Ala Gly Trp
 1 5

 <210> 141
 <211> 8
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 141
 Arg Ser Val Ser Ser Phe Gly Trp
 1 5

 <210> 142
 <211> 8
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 <220>
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 Trp residue

 <400> 142
 Leu Gly Thr Arg Lys Gly Gly Trp
 1 5

 <210> 143
 <211> 8
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<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 143

Lys Asp Pro Leu Phe Asn Gly Trp
1 5

<210> 144

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 144

Leu Ser Gln His Thr Asn Gly Trp
1 5

<210> 145

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 145

Asn Arg Leu Leu Leu Thr Gly Trp
1 5

<210> 146

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 146

Tyr Pro Leu Trp Val Ile Gly Trp
1 5

<210> 147

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 147
 Leu Leu Ile Ile Asp Arg Gly Trp
 1 5

<210> 148
 <211> 8
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 Trp residue

<400> 148
 Arg Val Ile Ser Leu Gln Gly Trp
 1 5

<210> 149
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 Trp residue

<400> 149
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 1 5

<210> 150
 <211> 8
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<220>
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 Trp residue

<400> 150
 Ser Ile Leu Arg Ser Thr Gly Trp
 1 5

<210> 151
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<220>
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 Trp residue

<400> 151
 Pro Gly Leu Val Trp Leu Gly Trp
 1 5

<210> 152
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 Trp residue

 <400> 152
 Val Lys Lys Leu Tyr Ile Gly Trp
 1 5

 <210> 153
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 153
 Asn Asn Arg Leu Leu Asp Gly Trp
 1 5

 <210> 154
 <211> 8
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 <220>
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 Trp residue

 <400> 154
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 1 5

 <210> 155
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 155
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 1 5

 <210> 156
 <211> 8
 <212> PRT
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<220>
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Trp residue

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1 5

<210> 157
<211> 8
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<220>
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Trp residue

<400> 157
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1 5

<210> 158
<211> 8
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<220>
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Trp residue

<400> 158
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1 5

<210> 159
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<212> PRT
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<220>
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Trp residue

<400> 159
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1 5

<210> 160
<211> 8
<212> PRT
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<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 160

Ser Val Leu Asp His Val Gly Trp
1 5

<210> 161

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 161

Asn Leu Leu Arg Arg Ala Gly Trp
1 5

<210> 162

<211> 8

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<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 162

Ser Gly Ile Ser Ala Trp Gly Trp
1 5

<210> 163

<211> 8

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<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 163

Phe Tyr Phe Trp Val Arg Gly Trp
1 5

<210> 164

<211> 8

<212> PRT

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<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 164

Lys Leu Phe Leu Pro Leu Gly Trp
1 5

<210> 165
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 <220>
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 Trp residue

 <400> 165
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 1 5

 <210> 166
 <211> 8
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 <220>
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 Trp residue

 <400> 166
 Thr His Ser Leu Ile Leu Gly Trp
 1 5

 <210> 167
 <211> 8
 <212> PRT
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 Trp residue

 <400> 167
 Leu Leu Leu Leu Ser Arg Gly Trp
 1 5

 <210> 168
 <211> 8
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 <220>
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 Trp residue

 <400> 168
 Leu Leu Arg Val Arg Ser Gly Trp
 1 5

 <210> 169
 <211> 8
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<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 169

Glu Arg Arg Ser Arg Gly Gly Trp
1 5

<210> 170

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 170

Arg Met Leu Gln Leu Ala Gly Trp
1 5

<210> 171

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 171

Arg Gly Trp Ala Asn Ser Gly Trp
1 5

<210> 172

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 172

Arg Pro Phe Tyr Ser Tyr Gly Trp
1 5

<210> 173

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 173
Ser Ser Ser Trp Asn Ala Gly Trp
1 5

<210> 174
<211> 8
<212> PRT
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<220>
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Trp residue

<400> 174
Leu Gly His Leu Glu Glu Gly Trp
1 5

<210> 175
<211> 8
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<220>
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Trp residue

<400> 175
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1 5

<210> 176
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<220>
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Trp residue

<400> 176
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1 5

<210> 177
<211> 7
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<220>
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Trp residue

<400> 177
Leu Arg Arg Trp Ser Leu Trp
1 5

<210> 178
 <211> 7
 <212> PRT
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 <220>
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 Trp residue

 <400> 178
 Lys Trp Val His Leu Phe Trp
 1 5

 <210> 179
 <211> 7
 <212> PRT
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 Trp residue

 <400> 179
 Asn Arg Leu Leu Leu Thr Trp
 1 5

 <210> 180
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 <212> PRT
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 Trp residue

 <400> 180
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 1 5

 <210> 181
 <211> 7
 <212> PRT
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 Trp residue

 <400> 181
 Asn Ala Leu Leu Leu Thr Trp
 1 5

 <210> 182
 <211> 7
 <212> PRT
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Trp residue

<400> 182
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1 5

<210> 183
<211> 7
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<220>
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Trp residue

<400> 183
Asn Leu Leu Arg Leu Thr Trp
1 5

<210> 184
<211> 7
<212> PRT
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<220>
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Trp residue

<400> 184
Asn Arg Leu Trp Leu Thr Trp
1 5

<210> 185
<211> 7
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<220>
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Trp residue

<400> 185
Asn Arg Leu Leu Leu Ala Trp
1 5

<210> 186
<211> 8
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<220>
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Trp residue

<400> 186
 Phe Tyr Gln Leu Ala Leu Thr Trp
 1 5

<210> 187
 <211> 8
 <212> PRT
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<220>
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 Trp residue

<400> 187
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 1 5

<210> 188
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<220>
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 Trp residue

<400> 188
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 1 5

<210> 189
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<220>
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 Trp residue

<400> 189
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 1 5

<210> 190
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<220>
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 Trp residue

<400> 190
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 1 5

<210> 191
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 <220>
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 Trp residue

 <400> 191
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 1 5

 <210> 192
 <211> 9
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 <220>
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 Trp residue

 <400> 192
 Arg Gly Tyr Val Tyr Gln Gly Leu Trp
 1 5

 <210> 193
 <211> 8
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 <220>
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 Trp residue

 <400> 193
 Tyr Thr Leu Val Gln Pro Leu Trp
 1 5

 <210> 194
 <211> 8
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 <220>
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 Trp residue

 <400> 194
 Thr Pro Asp Ile Thr Pro Lys Trp
 1 5

 <210> 195
 <211> 8
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<220>
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Trp residue

<400> 195
Thr Tyr Pro Asp Leu Arg Tyr Trp
1 5

<210> 196
<211> 8
<212> PRT
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<220>
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Trp residue

<400> 196
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1 5

<210> 197
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
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Trp residue

<400> 197
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1 5

<210> 198
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 198
Tyr Gln His Ala Val Gln Thr Trp
1 5

<210> 199
<211> 8
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<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 199
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1 5

<210> 200
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
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Trp residue

<400> 200
Ser Ser Phe Pro Pro Leu Asp Trp
1 5

<210> 201
<211> 8
<212> PRT
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<220>
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Trp residue

<400> 201
Met Ala Pro Ser Pro Pro His Trp
1 5

<210> 202
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 202
Ser Ser Phe Pro Asp Leu Leu Trp
1 5

<210> 203
<211> 8
<212> PRT
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<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 203
His Ser Tyr Asn Arg Leu Pro Trp
1 5

<210> 204
 <211> 8
 <212> PRT
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 <220>
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 Trp residue

 <400> 204
 His Leu Thr His Ser Gln Arg Trp
 1 5

 <210> 205
 <211> 8
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 <213> Artificial Sequence

 <220>
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 Trp residue

 <400> 205
 Gln Ala Ala Gln Ser Arg Ser Trp
 1 5

 <210> 206
 <211> 8
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 206
 Phe Ala Thr His His Ile Gly Trp
 1 5

 <210> 207
 <211> 8
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 <220>
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 Trp residue

 <400> 207
 Ser Met Pro Glu Pro Leu Ile Trp
 1 5

 <210> 208
 <211> 8
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<220>
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Trp residue

<400> 208
Ile Pro Arg Tyr His Leu Ile Trp
1 5

<210> 209
<211> 8
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<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

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1 5

<210> 210
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<220>
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Trp residue

<400> 210
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1 5

<210> 211
<211> 8
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<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 211
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1 5

<210> 212
<211> 8
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<213> Artificial Sequence

<220>
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Trp residue

<400> 212

Ala Ser Ala Gly Tyr Gln Ile Trp
1 5

<210> 213
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
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Trp residue

<400> 213
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1 5

<210> 214
<211> 8
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<220>
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Trp residue

<400> 214
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1 5

<210> 215
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 215
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1 5

<210> 216
<211> 8
<212> PRT
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<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 216
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1 5

<210> 217

<211> 8
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 217
 Gly Gln Trp Trp Ser Pro Asp Trp
 1 5

 <210> 218
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 218
 Gly Pro Pro His Gln Asp Ser Trp
 1 5

 <210> 219
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 219
 Asn Thr Leu Pro Ser Thr Ile Trp
 1 5

 <210> 220
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 220
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 1 5

 <210> 221
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 221

Tyr Gly Asn Pro Leu Gln Pro Trp
1 5

<210> 222

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 222

Phe His Trp Trp Trp Gln Pro Trp
1 5

<210> 223

<211> 8

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<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 223

Ile Thr Leu Lys Tyr Pro Leu Trp
1 5

<210> 224

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 224

Phe His Trp Pro Trp Leu Phe Trp
1 5

<210> 225

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 225

Thr Ala Gln Asp Ser Thr Gly Trp

1 5

<210> 226
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<220>
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 Trp residue

<400> 226
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 1 5

<210> 227
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<220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

<400> 227
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 1 5

<210> 228
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
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 Trp residue

<400> 228
 Glu Pro Phe Phe Arg Met Gln Trp
 1 5

<210> 229
 <211> 8
 <212> PRT
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<220>
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 Trp residue

<400> 229
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 1 5

<210> 230
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<212> PRT
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 <220>
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 Trp residue

 <400> 230
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 1 5

 <210> 231
 <211> 8
 <212> PRT
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 <220>
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 Trp residue

 <400> 231
 Gln Pro Ser His Leu Arg Trp Trp
 1 5

 <210> 232
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 232
 Ser Pro Ala Ser Pro Val Tyr Trp
 1 5

 <210> 233
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 233
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 1 5

 <210> 234
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 <220>
 <223> Heat shock protein binding domain with a terminal

Trp residue

<400> 234

His Pro Ser Asn Gln Ala Ser Trp
1 5

<210> 235

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 235

Asn Ser Ala Pro Arg Pro Val Trp
1 5

<210> 236

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 236

Gln Leu Trp Ser Ile Tyr Pro Trp
1 5

<210> 237

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 237

Ser Trp Pro Phe Phe Asp Leu Trp
1 5

<210> 238

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 238

Asp Thr Thr Leu Pro Leu His Trp
1 5

<210> 239
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 239
 Trp His Trp Gln Met Leu Trp Trp
 1 5

 <210> 240
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 240

 Asp Ser Phe Arg Thr Pro Val Trp
 1 5

 <210> 241
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 241
 Thr Ser Pro Leu Ser Leu Leu Trp
 1 5

 <210> 242
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 <220>
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 Trp residue

 <400> 242
 Ala Tyr Asn Tyr Val Ser Asp Trp
 1 5

 <210> 243
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<212> PRT
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 Trp residue

 <400> 243
 Arg Pro Leu His Asp Pro Met Trp
 1 5

 <210> 244
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 <220>
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 Trp residue

 <400> 244
 Trp Pro Ser Thr Thr Leu Phe Trp
 1 5

 <210> 245
 <211> 8
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 <220>
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 Trp residue

 <400> 245
 Ala Thr Leu Glu Pro Val Arg Trp
 1 5

 <210> 246
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 246
 Ser Met Thr Val Leu Arg Pro Trp
 1 5

 <210> 247
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with a terminal

Trp residue

<400> 247

Gln Ile Gly Ala Pro Ser Trp Trp
1 5

<210> 248

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal

Trp residue

<400> 248

Ala Pro Asp Leu Tyr Val Pro Trp
1 5

<210> 249

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 249

Arg Met Pro Pro Leu Leu Pro Trp
1 5

<210> 250

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 250

Ala Lys Ala Thr Pro Glu His Trp
1 5

<210> 251

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 251

Thr Pro Pro Leu Arg Ile Asn Trp

1 5

<210> 252
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<220>
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 Trp residue

<400> 252
 Leu Pro Ile His Ala Pro His Trp
 1 5

<210> 253
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 Trp residue

<400> 253
 Asp Leu Asn Ala Tyr Thr His Trp
 1 5

<210> 254
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<220>
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 Trp residue

<400> 254
 Val Thr Leu Pro Asn Phe His Trp
 1 5

<210> 255
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 Trp residue

<400> 255
 Asn Ser Arg Leu Pro Thr Leu Trp
 1 5

<210> 256
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<212> PRT
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 <223> Heat shock protein binding domain with a terminal
 Trp residue

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 Tyr Pro His Pro Ser Arg Ser Trp
 1 5

 <210> 257
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 <220>
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 Trp residue

 <400> 257
 Gly Thr Ala His Phe Met Tyr Trp
 1 5

 <210> 258
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 <220>
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 Trp residue

 <400> 258
 Tyr Ser Leu Leu Pro Thr Arg Trp
 1 5

 <210> 259
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 259
 Leu Pro Arg Arg Thr Leu Leu Trp
 1 5

 <210> 260
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 <220>
 <223> Heat shock protein binding domain with a terminal

Trp residue

<400> 260

Thr Ser Thr Leu Leu Trp Lys Trp
1 5

<210> 261

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 261

Thr Ser Asp Met Lys Pro His Trp
1 5

<210> 262

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 262

Thr Ser Ser Tyr Leu Ala Leu Trp
1 5

<210> 263

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 263

Asn Leu Tyr Gly Pro His Asp Trp
1 5

<210> 264

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 264

Leu Glu Thr Tyr Thr Ala Ser Trp
1 5

<210> 265
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 265
 Ala Tyr Lys Ser Leu Thr Gln Trp
 1 5

 <210> 266
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 266
 Ser Thr Ser Val Tyr Ser Ser Trp
 1 5

 <210> 267
 <211> 8
 <212> PRT
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 <220>
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 Trp residue

 <400> 267
 Glu Gly Pro Leu Arg Ser Pro Trp
 1 5

 <210> 268
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 268
 Thr Thr Tyr His Ala Leu Gly Trp
 1 5

 <210> 269
 <211> 8
 <212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 269

Val Ser Ile Gly His Pro Ser Trp
1 5

<210> 270

<211> 8

<212> PRT

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<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 270

Thr His Ser His Arg Pro Ser Trp
1 5

<210> 271

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 271

Ile Thr Asn Pro Leu Thr Thr Trp
1 5

<210> 272

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 272

Ser Ile Gln Ala His His Ser Trp
1 5

<210> 273

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 273
 Leu Asn Trp Pro Arg Val Leu Trp
 1 5

<210> 274
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<220>
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 Trp residue

<400> 274
 Tyr Tyr Tyr Ala Pro Pro Pro Trp
 1 5

<210> 275
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<220>
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 Trp residue

<400> 275
 Ser Leu Trp Thr Arg Leu Pro Trp
 1 5

<210> 276
 <211> 8
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<220>
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 Trp residue

<400> 276
 Asn Val Tyr His Ser Ser Leu Trp
 1 5

<210> 277
 <211> 8
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<220>
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 Trp residue

<400> 277
 Asn Ser Pro His Pro Pro Thr Trp
 1 5

<210> 278
 <211> 8
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 <220>
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 Trp residue

 <400> 278
 Val Pro Ala Lys Pro Arg His Trp
 1 5

 <210> 279
 <211> 8
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 279
 His Asn Leu His Pro Asn Arg Trp
 1 5

 <210> 280
 <211> 8
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 280
 Tyr Thr Thr His Arg Trp Leu Trp
 1 5

 <210> 281
 <211> 8
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 <220>
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 Trp residue

 <400> 281
 Ala Val Thr Ala Ala Ile Val Trp
 1 5

 <210> 282
 <211> 8
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<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 282
Thr Leu Met His Asp Arg Val Trp
1 5

<210> 283
<211> 8
<212> PRT
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<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 283
Thr Pro Leu Lys Val Pro Tyr Trp
1 5

<210> 284
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 284
Phe Thr Asn Gln Gln Tyr His Trp
1 5

<210> 285
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 285
Ser His Val Pro Ser Met Ala Trp
1 5

<210> 286
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 286
His Thr Thr Val Tyr Gly Ala Trp
1 5

<210> 287
<211> 8
<212> PRT
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<220>
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Trp residue

<400> 287
Thr Glu Thr Pro Tyr Pro Thr Trp
1 5

<210> 288
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 288
Leu Thr Thr Pro Phe Ser Ser Trp
1 5

<210> 289
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 289
Gly Val Pro Leu Thr Met Asp Trp
1 5

<210> 290
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 290
Lys Leu Pro Thr Val Leu Arg Trp
1 5

<210> 291
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 291
 Cys Arg Phe His Gly Asn Arg Trp
 1 5

 <210> 292
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 292
 Tyr Thr Arg Asp Phe Glu Ala Trp
 1 5

 <210> 293
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 293
 Ser Ser Ala Ala Gly Pro Arg Trp
 1 5

 <210> 294
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 294
 Ser Leu Ile Gln Tyr Ser Arg Trp
 1 5

 <210> 295
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<221> VARIANT
<222> 7
<223> Xaa = any amino acid

<400> 295
Asp Ala Leu Met Trp Pro Xaa Trp
1 5

<210> 296
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<221> VARIANT
<222> 3
<223> Xaa = any amino acid

<400> 296
Ser Ser Xaa Ser Leu Tyr Ile Trp
1 5

<210> 297
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 297
Phe Asn Thr Ser Thr Arg Thr Trp
1 5

<210> 298
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 298
Thr Val Gln His Val Ala Phe Trp
1 5

<210> 299
<211> 8

<212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 299
 Asp Tyr Ser Phe Pro Pro Leu Trp
 1 5

 <210> 300
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 300
 Val Gly Ser Met Glu Ser Leu Trp
 1 5

 <210> 301
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <221> VARIANT
 <222> 2, 6
 <223> Xaa = any amino acid

 <400> 301
 Phe Xaa Pro Met Ile Xaa Ser Trp
 1 5

 <210> 302
 <211> 8
 <212> PRT
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 <220>
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 Trp residue

 <400> 302
 Ala Pro Pro Arg Val Thr Met Trp
 1 5

 <210> 303
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<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 303

Ile Ala Thr Lys Thr Pro Lys Trp
1 5

<210> 304

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 304

Lys Pro Pro Leu Phe Gln Ile Trp
1 5

<210> 305

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 305

Tyr His Thr Ala His Asn Met Trp
1 5

<210> 306

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 306

Ser Tyr Ile Gln Ala Thr His Trp
1 5

<210> 307

<211> 8

<212> PRT

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<220>

<223> Heat shock protein binding domain with a terminal

Trp residue

<400> 307

Ser Ser Phe Ala Thr Phe Leu Trp
1 5

<210> 308

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 308

Thr Thr Pro Pro Asn Phe Ala Trp
1 5

<210> 309

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 309

Ile Ser Leu Asp Pro Arg Met Trp
1 5

<210> 310

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 310

Ser Leu Pro Leu Phe Gly Ala Trp
1 5

<210> 311

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 311

Asn Leu Leu Lys Thr Thr Leu Trp
1 5

<210> 312
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 <220>
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 Trp residue

 <400> 312
 Asp Gln Asn Leu Pro Arg Arg Trp
 1 5

 <210> 313
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 313
 Ser His Phe Glu Gln Leu Leu Trp
 1 5

 <210> 314
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 314
 Thr Pro Gln Leu His His Gly Trp
 1 5

 <210> 315
 <211> 8

 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 315
 Ala Pro Leu Asp Arg Ile Thr Trp
 1 5

 <210> 316
 <211> 8

<212> PRT
 <213> Artificial Sequence

 <220>
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 Trp residue

 <400> 316
 Phe Ala Pro Leu Ile Ala His Trp
 1 5

 <210> 317
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 317
 Ser Trp Ile Gln Thr Phe Met Trp
 1 5

 <210> 318
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 318
 Asn Thr Trp Pro His Met Tyr Trp
 1 5

 <210> 319
 <211> 8
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 <220>
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 Trp residue

 <400> 319
 Glu Pro Leu Pro Thr Thr Leu Trp
 1 5

 <210> 320
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
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Trp residue

<400> 320

His Gly Pro His Leu Phe Asn Trp
1 5

<210> 321

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 321

Tyr Leu Asn Ser Thr Leu Ala Trp
1 5

<210> 322

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 322

His Leu His Ser Pro Ser Gly Trp
1 5

<210> 323

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 323

Thr Leu Pro His Arg Leu Asn Trp
1 5

<210> 324

<211> 8

<212> PRT

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<220>

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Trp residue

<400> 324

Ser Ser Pro Arg Glu Val His Trp
1 5

<210> 325
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 <220>
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 Trp residue

 <400> 325
 Asn Gln Val Asp Thr Ala Arg Trp
 1 5

 <210> 326
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 Trp residue

 <400> 326
 Tyr Pro Thr Pro Leu Leu Thr Trp
 1 5

 <210> 327
 <211> 8
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 Trp residue

 <400> 327
 His Pro Ala Ala Phe Pro Trp Trp
 1 5

 <210> 328
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 Trp residue

 <400> 328
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 1 5

 <210> 329
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<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 329

Leu Glu Thr Tyr Thr Ala Ser Trp
1 5

<210> 330

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 330

Lys Tyr Val Pro Leu Pro Pro Trp
1 5

<210> 331

<211> 8

<212> PRT

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<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 331

Ala Pro Leu Ala Leu His Ala Trp
1 5

<210> 332

<211> 8

<212> PRT

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<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 332

Tyr Glu Ser Leu Leu Thr Lys Trp
1 5

<210> 333

<211> 8

<212> PRT

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<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 333
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 1 5

<210> 334
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<220>
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 Trp residue

<400> 334
 Gly Leu Ala Thr Val Lys Ser Trp
 1 5

<210> 335
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 Trp residue

<400> 335
 Gly Ala Thr Ser Phe Gly Leu Trp
 1 5

<210> 336
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 Trp residue

<400> 336
 Lys Pro Pro Gly Pro Val Ser Trp
 1 5

<210> 337
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 <213> Artificial Sequence

<220>
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 Trp residue

<400> 337
 Thr Leu Tyr Val Ser Gly Asn Trp
 1 5

<210> 338
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 338
 His Ala Pro Phe Lys Ser Gln Trp
 1 5

 <210> 339
 <211> 8
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 Trp residue

 <400> 339
 Val Ala Phe Thr Arg Leu Pro Trp
 1 5

 <210> 340
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 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 340
 Leu Pro Thr Arg Thr Pro Ala Trp
 1 5

 <210> 341
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 Trp residue

 <400> 341
 Ala Ser Phe Asp Leu Leu Ile Trp
 1 5

 <210> 342
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Trp residue

<400> 342

Arg Met Asn Thr Glu Pro Pro Trp
1 5

<210> 343

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<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 343

Lys Met Thr Pro Leu Thr Thr Trp
1 5

<210> 344

<211> 8

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<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 344

Ala Asn Ala Thr Pro Leu Leu Trp
1 5

<210> 345

<211> 8

<212> PRT

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Trp residue

<400> 345

Thr Ile Trp Pro Pro Pro Val Trp
1 5

<210> 346

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 346
 Gln Thr Lys Val Met Thr Thr Trp
 1 5

<210> 347
 <211> 8
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 <213> Artificial Sequence

<220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

<400> 347
 Asn His Ala Val Phe Ala Ser Trp
 1 5

<210> 348
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

<221> VARIANT
 <222> 5
 <223> Xaa = any amino acid

<400> 348
 Leu His Ala Ala Xaa Thr Ser Trp
 1 5

<210> 349
 <211> 8
 <212> PRT
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<220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

<400> 349
 Thr Trp Gln Pro Tyr Phe His Trp
 1 5

<210> 350
 <211> 8
 <212> PRT
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<220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

<400> 350
Ala Pro Leu Ala Leu His Ala Trp
1 5

<210> 351
<211> 8
<212> PRT
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<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 351
Thr Ala His Asp Leu Thr Val Trp
1 5

<210> 352
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 352
Asn Met Thr Asn Met Leu Thr Trp
1 5

<210> 353
<211> 8
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<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 353
Gly Ser Gly Leu Ser Gln Asp Trp
1 5

<210> 354
<211> 8
<212> PRT
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<220>
<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 354
Thr Pro Ile Lys Thr Ile Tyr Trp
1 5

<210> 355
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 355
 Ser His Leu Tyr Arg Ser Ser Trp
 1 5

 <210> 356
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with a terminal
 Trp residue

 <400> 356
 His Gly Gln Ala Trp Gln Phe Trp
 1 5

 <210> 357
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain

 <400> 357
 Ser Ile Ile Asn Phe Glu Lys Leu
 1 5

 <210> 358
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain

 <400> 358
 His Trp Asp Phe Ala Trp Pro Trp
 1 5

 <210> 359
 <211> 8
 <212> PRT
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 <220>

<223> Heat shock protein binding domain

<400> 359

Asn Leu Leu Arg Leu Thr Gly Trp

1

5

<210> 360

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain

<400> 360

Phe Tyr Gln Leu Ala Leu Thr Trp

1

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<210> 361

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain

<400> 361

Arg Lys Leu Phe Phe Asn Leu Arg Trp

1

5

<210> 362

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain

<400> 362

Ala Leu Phe Asp Ile Glu Ser Lys Val

1

5

<210> 363

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain

<400> 363

Ile Met Asp Gln Val Pro Phe Ser Val

1

5

<210> 364

<211> 9

<212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain

 <400> 364
 Tyr Met Asp Gly Thr Met Ser Gln Val
 1 5

 <210> 365
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain

 <400> 365
 Thr Leu Gly Ile Val Cys Pro Ile
 1 5

 <210> 366
 <211> 10
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain

 <400> 366
 Tyr Met Leu Asp Leu Gln Pro Glu Thr Thr
 1 5 10

 <210> 367
 <211> 19
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Hybrid antigen

 <400> 367
 Ser Ile Ile Asn Phe Glu Lys Leu Gly Ser Gly Asn Leu Leu Arg Leu
 1 5 10 15
 Thr Gly Trp

 <210> 368
 <211> 19
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Hybrid antigen

 <400> 368

Ser Ile Ile Asn Phe Glu Lys Leu Gly Ser Gly His Trp Asp Phe Ala

1 5 10 15
Trp Pro Trp

<210> 369
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 369
Ala Leu Phe Asp Ile Glu Ser Lys Val Gly Ser Gly His Trp Asp Phe
1 5 10 15
Ala Trp Pro Trp
20

<210> 370
<211> 8
<212> PRT
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<220>
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<400> 370
Arg Gly Tyr Val Tyr Gln Gly Leu
1 5

<210> 371
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain

<400> 371
Ile Met Asp Gln Val Pro Phe Ser Val Gly Ser Gly His Trp Asp Phe
1 5 10 15
Ala Trp Pro Trp
20

<210> 372
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 372
Ile Met Asp Gln Val Pro Phe Ser Val Gly Ser Gly Asn Leu Leu Arg
1 5 10 15

Leu Thr Gly Trp
20

<210> 373
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 373
Tyr Met Asp Gly Thr Met Ser Gln Val Gly Ser Gly His Trp Asp Phe
1 5 10 15
Ala Trp Pro Trp
20

<210> 374
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 374
His Trp Asp Phe Ala Trp Pro Trp Gly Ser Gly Tyr Met Asp Gly Thr
1 5 10 15
Met Ser Gln Val
20

<210> 375
<211> 23
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 375
Tyr Met Asp Gly Thr Met Ser Gln Val Gly Ser Gly Gly Ser Gly Asn
1 5 10 15
Leu Leu Arg Leu Thr Gly Trp
20

<210> 376
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 376
Thr Leu Gly Ile Val Cys Pro Ile Gly Ser Gly His Trp Asp Phe Ala
1 5 10 15
Trp Pro Trp

<210> 377
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 377
Thr Leu Gly Ile Val Cys Pro Ile Gly Ser Gly Gly Asn Leu Leu Arg
1 5 10 15
Leu Thr Gly Trp
20

<210> 378
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 378
Tyr Met Leu Asp Leu Gln Pro Glu Thr Thr Gly Ser Gly His Trp Asp
1 5 10 15
Phe Ala Trp Pro Trp
20

<210> 379
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
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<400> 379
His Trp Asp Phe Ala Trp Pro Trp Gly Ser Gly Ser Ile Ile Asn Phe
1 5 10 15
Glu Lys Leu

<210> 380
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
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<400> 380
Ser Ile Ile Asn Phe Glu Lys Leu Gly Ser Gly Asn Leu Leu Arg Leu
1 5 10 15
Thr Gly Trp

<210> 381
 <211> 19
 <212> PRT
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 <220>
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 <400> 381
 Ser Ile Ile Asn Phe Glu Lys Leu Gly Ser Gly Phe Tyr Gln Leu Ala
 1 5 10 15
 Leu Thr Trp

<210> 382
 <211> 20
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 <220>
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 <400> 382
 Ser Ile Ile Asn Phe Glu Lys Leu Gly Ser Gly Arg Lys Leu Phe Phe
 1 5 10 15
 Asn Leu Arg Trp
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<210> 383
 <211> 19
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 <220>
 <223> Heat shock protein binding domain

 <400> 383
 Asn Leu Leu Arg Leu Thr Gly Trp Gly Ser Gly Ser Ile Ile Asn Phe
 1 5 10 15
 Glu Lys Leu

<210> 384
 <211> 20
 <212> PRT
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 <220>
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 <400> 384
 Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Ile Ile Asn
 1 5 10 15
 Phe Glu Lys Leu
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<210> 385
 <211> 18
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain

 <400> 385
 Asn Leu Leu Arg Leu Thr Gly Trp Arg Lys Ser Ile Ile Asn Phe Glu
 1 5 10 15
 Lys Leu

<210> 386
 <211> 19
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain

 <400> 386
 Asn Leu Leu Arg Leu Thr Gly Trp Gly Ser Gly Arg Gly Tyr Val Tyr
 1 5 10 15
 Gln Gly Leu

<210> 387
 <211> 20
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain

 <400> 387
 Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Arg Gly Tyr Val
 1 5 10 15
 Tyr Gln Gly Leu
 20

<210> 388
 <211> 18
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain

 <400> 388
 Asn Leu Leu Arg Leu Thr Gly Trp Arg Lys Arg Gly Tyr Val Tyr Gln
 1 5 10 15
 Gly Leu

<210> 389
 <211> 10
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain

 <400> 389
 Glu Leu Ala Gly Ile Gly Ile Leu Thr Val
 1 5 10

 <210> 390
 <211> 9
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain

 <400> 390
 Ser Leu Leu Met Trp Ile Thr Gln Val
 1 5

 <210> 391
 <211> 9
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain

 <400> 391
 Ser Val Tyr Asp Phe Phe Val Trp Leu
 1 5

 <210> 392
 <211> 9
 <212> PRT
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 <220>
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 <400> 392
 Gly Leu Tyr Asp Gly Met Glu His Leu
 1 5

 <210> 393
 <211> 9
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain

 <400> 393

Tyr Leu Glu Pro Gly Pro Val Thr Val
1 5

<210> 394
<211> 9
<212> PRT
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<220>
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<400> 394
Lys Ala Ser Glu Lys Ile Phe Tyr Val
1 5

<210> 395
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 395
Glu Leu Ala Gly Ile Gly Ile Leu Thr Val Gly Ser Gly Asn Leu Leu
1 5 10 15
Arg Leu Thr Gly Trp
20

<210> 396
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 396
Ser Leu Leu Met Trp Ile Thr Gln Val Gly Ser Gly Asn Leu Leu Arg
1 5 10 15
Leu Thr Gly Trp
20

<210> 397
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
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<400> 397
Ser Val Tyr Asp Phe Phe Val Trp Leu Gly Ser Gly Asn Leu Leu Arg
1 5 10 15
Leu Thr Gly Trp
20

<210> 398
 <211> 20
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Hybrid antigen

 <400> 398
 Gly Leu Tyr Asp Gly Met Glu His Leu Gly Ser Gly Asn Leu Leu Arg
 1 5 10 15
 Leu Thr Gly Trp
 20

<210> 399
 <211> 20
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Hybrid antigen

 <400> 399
 Tyr Leu Glu Pro Gly Pro Val Thr Val Gly Ser Gly Asn Leu Leu Arg
 1 5 10 15
 Leu Thr Gly Trp
 20

<210> 400
 <211> 20
 <212> PRT
 <213> Artificial Sequence

 <220>
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 <400> 400
 Lys Ala Ser Glu Lys Ile Phe Tyr Val Gly Ser Gly Asn Leu Leu Arg
 1 5 10 15
 Leu Thr Gly Trp
 20

<210> 401
 <211> 9
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain

 <400> 401
 Ala Leu Lys His Arg Ala Tyr Glu Leu
 1 5

<210> 402

<211> 9
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain

 <400> 402
 Ile Leu Lys Glu Pro Val His Gly Val
 1 5

 <210> 403
 <211> 9
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain

 <400> 403
 Ser Leu Phe Asn Thr Val Ala Thr Leu
 1 5

 <210> 404
 <211> 11
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain

 <400> 404
 Val Leu Asp Val Gly Asp Ala Tyr Phe Ser Val
 1 5 10

 <210> 405
 <211> 9
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain

 <400> 405
 Val Ile Tyr Gln Tyr Met Asp Asp Leu
 1 5

 <210> 406
 <211> 9
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain

 <400> 406
 Ser Leu Tyr Asn Thr Val Ala Thr Leu

1 5

<210> 407
 <211> 9
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heat shock protein binding domain

<400> 407
 Ala Ile Ile Arg Ile Leu Gln Gln Leu
 1 5

<210> 408
 <211> 9
 <212> PRT
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<220>
 <223> Heat shock protein binding domain

<400> 408
 Ala Phe His His Val Ala Arg Glu Leu
 1 5

<210> 409
 <211> 20
 <212> PRT
 <213> Artificial Sequence

<220>
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<400> 409
 Ala Leu Lys His Arg Ala Tyr Glu Leu Gly Ser Gly Asn Leu Leu Arg
 1 5 10 15
 Leu Thr Gly Trp
 20

<210> 410
 <211> 20
 <212> PRT
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<220>
 <223> Hybrid antigen

<400> 410
 Ile Leu Lys Glu Pro Val His Gly Val Gly Ser Gly Asn Leu Leu Arg
 1 5 10 15
 Leu Thr Gly Trp
 20

<210> 411
 <211> 20

<212> PRT
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<220>
<223> Hybrid antigen

<400> 411
Ser Leu Phe Asn Thr Val Ala Thr Leu Gly Ser Gly Asn Leu Leu Arg
1 5 10 15
Leu Thr Gly Trp
20

<210> 412
<211> 22
<212> PRT
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<220>
<223> Hybrid antigen

<400> 412
Val Leu Asp Val Gly Asp Ala Tyr Phe Ser Val Gly Ser Gly Asn Leu
1 5 10 15
Leu Arg Leu Thr Gly Trp
20

<210> 413
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
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<400> 413
Val Ile Tyr Gln Tyr Met Asp Asp Leu Gly Ser Gly Asn Leu Leu Arg
1 5 10 15
Leu Thr Gly Trp
20

<210> 414
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 414
Ser Leu Tyr Asn Thr Val Ala Thr Leu Gly Ser Gly Asn Leu Leu Arg
1 5 10 15
Leu Thr Gly Trp
20

<210> 415
<211> 20
<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 415

Ala Ile Ile Arg Ile Leu Gln Gln Leu Gly Ser Gly Asn Leu Leu Arg
1 5 10 15
Leu Thr Gly Trp
20

<210> 416

<211> 20

<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 416

Ala Phe His His Val Ala Arg Glu Leu Gly Ser Gly Asn Leu Leu Arg
1 5 10 15
Leu Thr Gly Trp
20

<210> 417

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 417

Asn Leu Leu Arg Leu Thr Gly Trp
1 5

<210> 418

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 418

Phe Tyr Gln Leu Ala Leu Tyr Trp
1 5

<210> 419

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain with a terminal

Trp residue

<400> 419

Arg Lys Leu Phe Phe Asn Leu Arg Trp
1 5